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(54) Title: ANALOGS OF PEPTIDE YY AND USES 7 (57) Abstract The invention provides analogs of PYY. The invention activities such as cell proliferation, nutrient transport, lips	ention al	provides compositions and methods useful for controlling	biologica

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ANALOGS OF PEPTIDE YY AND USES THEREOF Statement as To Federally Sponsored Research

This invention was made in part with Government funding and the Government therefore has certain rights in the invention.

Background of the Invention

This invention relates to peptide derivatives which are useful as therapeutic agents in the treatment of gastroenterological disorders.

Peptide YY (PYY) is a 36-residue peptide amide isolated originally from porcine intestine, and localized in the endocrine cells of the gastrointestinal tract and pancreas (Tatemoto et al. Proc. Natl. Acad. Sci. 79:2514, 15 1982). Peptide YY has N-terminal and C-terminal tyrosine amides; accordingly, these two tyrosines give PYY its name (Y represents the amino acid tyrosine in the peptide nomenclature). In addition PYY shares a number of central and peripheral regulatory roles with its 20 homologous peptide neuropeptide Y (NPY), which was originally isolated from porcine brain (Tatemoto, Proc. Natl. Acad. Sci. 79:5485, 1982). In contrast with the cellular location of PYY, NPY is present in submucous and myenteric neurons which innervate the mucosal and smooth 25 muscle layers, respectively (Ekblad et al. Neuroscience 20:169, 1987). Both PYY and NPY are believed to inhibit gut motility and blood flow (Laburthe, Trends Endocrinol. Metab. 1:168, 1990), and they are also thought to attenuate basal (Cox et al. Br. J. Pharmacol. 101:247, 30 1990; Cox et al. J. Physiol. 398:65, 1988; Cox et al. Peptides 12:323, 1991; Friel et al. Br. J. Pharmacol. 88:425, 1986) and secretagogue-induced intestinal secretion in rats (Lundberg et al. Proc. Natl. Acad. Sci USA 79:4471, 1982; Playford et al. Lancet 335:1555, 1990) 35 and humans (Playford et al. supra), as well as stimulate

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net absorption (MacFadyen et al. Neuropeptides 7:219, 1986). Furthermore, plasma PYY levels have been reported to be elevated in several diseases that cause diarrhea (Adrian et al. Gastroenterology 89:1070, 1985). Taken together, these observations suggest that PYY and NPY are released into the circulation after a meal (Adrian et al. Gastroenterology 89:1070, 1985; Balasubramaniam et al. Neuropeptides 14:209, 1989), and thus may play a physiological role in regulating intestinal secretion and absorption, serving as natural inhibitors of diarrhea.

A high affinity PYY receptor system which exhibits a slightly higher affinity for PYY than NPY has been characterized in rat intestinal epithelia (Laburthe et al. Endocrinology 118:1910, 1986; Laburthe, Trends

15 Endocrinol. Metab. supra) and shown to be negatively coupled to adenylate cyclase (Servin et al. Endocrinology 124:692, 1989). Consistently, PYY exhibited greater antisecretory potency than NPY in voltage clamped preparations of rat small intestine (Cox et al. J.

20 Physiol. supra), while
C-terminal fragments of NPY were found to be less
effective in their antisecretory potency than PYY (Cox et
al. Br. J. Pharmacol. supra). Structure-activity studies
using several partial sequences have led to the
25 identification of PYY(22-36) as the active site for
interacting with intestinal PYY receptors (Balsubramaniam
et al. Pept. Res. 1:32, 1988).

In addition, PYY has been implicated in a number of physiological activities including nutrient uptake (see, e.g., Bilcheik et al. Digestive Disease Week 506:623, 1993), cell proliferation (see, e.g., Laburthe, Trends Endocrinol. Metab. 1:168, 1990; Voisin et al. J. Biol. Chem, 1993), lipolysis (see, e.g., Valet et al., J. Clin. Invest. 85:291, 1990), and vasoconstriction

(see, e.g., Lundberg et al., Proc. Natl. Acad. SCi., USA 79: 4471, 1982).

The amino acid sequences of porcine and human PYY are as follows:

5 porcine PYY YPAKPEAPGEDASPEELSRYYASLRHYLNLVTRQRY (SEQ. ID. NO. 1)

human PYY YPIKPEAPGEDASPEELNRYYASLRHYLNLVTRQRY (SEQ. ID. NO. 2)

The amino acid sequence for dog PYY and rat is the same 10 as porcine PYY.

Summary of the Invention

In one aspect, the present invention features novel analogs of peptide YY of the formula:

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$$R_1$$
 R_2 R_2 R_2 R_3 R_2 R_3 R_4 R_2 R_4 (Formula I)

wherein

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X is a chain of 0-5 amino acids, inclusive, the N-20 terminal one of which is bonded to R_1 and R_2 ;

Y is a chain of 0-4 amino acids, inclusive, the C-terminal one of which is bonded to R_3 and R_4 ;

 R_1 is H, C_1 - C_{12} alkyl (e.g., methyl), C_6 - C_{18} aryl (e.g., phenyl, naphthaleneacetyl), C_1 - C_{12} acyl (e.g., formyl, acetyl, and myristoyl), C_7 - C_{18} aralkyl (e.g., benzyl), or C_7 - C_{18} alkaryl (e.g., p-methylphenyl);

 R_2 is H, C_1-C_{12} alkyl (e.g., methyl), C_6-C_{18} aryl (e.g., phenyl,naphthaleneacetyl), C_1-C_{12} acyl (e.g., formyl, acetyl, and myristoyl), C_7-C_{18} aralkyl (e.g.,benzyl), or C_7-C_{18} alkaryl (e.g., p-methylphenyl);

A²² is an aromatic amino acid, Ala, Aib, Anb, N-Me-Ala, or is deleted;

A²³ is Ser, Thr, Ala, Aib, N-Me-Ser, N-Me-Thr, N Ala, or is deleted; Me-A²⁴ is Leu, Ile, Val, Trp, Gly, Nle, Nva, Aib, Anb, N-Me-Leu, or is deleted; 5 A²⁵ is Arg, Lys, homo-Arg, diethyl-homo-Arg, Lys-NH-R (where R is H, a branched or straight ϵ $\dot{-}$ chain C_1-C_{10} alkyl group, or an aryl group), Orn, or is deleted; A^{26} is Ala, His, Thr, 3-Me-His, 1-Me-His, β -10 pyrozolylalanine, N-Me-His, Arg, Lys, homo-Arg, diethyl-homo-Arg, Lys- ϵ -NH-R (where R is H, a branched or straight chain C₁-C₁₀ alkyl group, or an aryl group), Orn, or is deleted; A²⁷ is an aromatic amino acid other than Tyr; 15 A²⁸ is Leu, Ile, Val, Trp, Nle, Nva, Aib, Anb, or N-Me-Leu; A²⁹ is Asn, Ala, Gln, Gly, Trp, or N-Me-Asn; A³⁰ is Leu, Ile, Val, Trp, Nle, Nva, Aib, Anb, or N-Me-Leu; A³¹ is Val, Leu, Ile, Trp, Nle, Nva, Aib, Anb, or N-Me-Val; A³² is Thr, Ser, N-Me-Ser, N-Me-Thr, or D-Trp; R_3 is H, C_1-C_{12} alkyl (e.g., methyl), C_6-C_{18} aryl (e.g., phenyl, naphthaleneacetyl), C_1-C_{12} 25 acyl (e.g., formyl, acetyl, and myristoyl), C_7-C_{18} aralkyl (e.g., benzyl), or C_7-C_{18} alkaryl (e.g., p-methylphenyl); and R_4 is H, C_1 - C_{12} alkyl (e.g., methyl), C_6 - C_{18} aryl (e.g., phenyl, naphthaleneacetyl), C₁-C₁₂ 30 acyl (e.g., formyl, acetyl, and myristoyl), C_7-C_{18} aralkyl (e.g., benzyl), or C_7-C_{18} alkaryl (e.g., p-methylphenyl), or a pharmaceutically acceptable salt thereof.

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In preferred embodiments, A^{27} is Phe, Nal, Bip, Pcp, Tic, Trp, Bth, Thi, or Dip.

In preferred embodiments X is $\mathtt{A}^{17}\text{-}\mathtt{A}^{18}\text{-}\mathtt{A}^{19}\text{-}\mathtt{A}^{20}\text{-}\mathtt{A}^{21}$ wherein

5 A¹⁷ is Cys, Leu, Ile, Val, Nle, Nva, Aib, Anb, or N-Me-Leu;

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A¹⁸ is Cys, Ser, Thr, N-Me-Ser, or N-Me-Thr;

 A^{19} is Arg, Lys, homo-Arg, diethyl-homo-Arg, Lys-NH-R (where R is H, a branched or straight chain C_1 - C_{10} alkyl group, or C_6 - C_{18} aryl group), Cys, or Orn;

A²⁰ is an aromatic amino acid, or Cys; and

 A^{21} is an aromatic amino acid, Cys, or a pharmaceutically acceptable salt thereof. In yet other preferred embodiments, Y is A^{33} - A^{34} - A^{35} - A^{36} wherein

 A^{33} is Arg, Lys, homo-Arg, diethyl-homo-Arg, Lys-NH-R (where R is H, a branched or straight chain C_1 - C_{10} alkyl group, or an aryl group), Cys, or Orn;

A³⁴ is Cys, Gln, Asn, Ala, Gly, N-Me-Gln, Aib, or Anb:

 A^{35} is Arg, Lys, homo-Arg, diethyl-homo-Arg, Lys- ϵ - NH-R(where R is H, a branched or straight chain C_1 - C_{10} alkyl group, or an aryl group), Cys, or Orn; and

A³⁶ is an aromatic amino acid, Cys or a pharmaceutically acceptable salt thereof.

Preferably, the compound has the formula: N-α-Ac-30 Ala-Ser-Leu-Arg-His-Phe-Leu-Asn-Leu-Val-Thr-Arg-Gln-Arg-Tyr-NH₂ (SEQ. ID. NO. 3), H-Ala-Ser-Leu-Arg-His-Phe-Leu-Asn-Leu-Val-Thr-Arg-Gln-Arg-Tyr-NH₂ (SEQ. ID. NO. 4), N-α-Ac-Ala-Ser-Leu-Arg-His-Trp-Leu-Asn-Leu-Val-Thr-Arg-Gln-Arg-Tyr-NH₂ (SEQ. ID. NO. 5), N-α-Ac-Ala-Ser-Leu-Arg-His-35 Thi-Leu-Asn-Leu-Val-Thr-Arg-Gln-Arg-Tyr-NH₂ (SEQ. ID. NO.

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6), $N-\alpha-Ac-Tyr-Ser-Leu-Arg-His-Phe-Leu-Asn-Leu-Val-Thr-Arg-Gln-Arg-Tyr-NH₂ (SEQ. ID. NO. 7) or a pharmaceutically acceptable salt thereof.$

In another aspect the invention features novel analogs of peptide YY of the formula:

$$R_1$$
 R_3 / $R_2-A^{25}-A^{26}-A^{27}-A^{28}-A^{29}-A^{30}-A^{31}-A^{32}-Y-R_4$ (Formula II)

wherein

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the N-terminal amino acid is bonded to R_1 and R_2 ; Y is a chain of 0-4 amino acids, inclusive the C-terminal one of which is bonded to R_3 and R_4 ;

 R_1 is H, C_1 - C_{12} alkyl (e.g., methyl), C_6 - C_{18} aryl (e.g., phenyl, napthaleneacetyl), C_1 - C_{12} acyl (e.g., formyl, acetyl, and myristoyl), C_7 - C_{18} aralkyl (e.g., benzyl), or C_7 - C_{18} alkaryl (e.g., p-methylphenyl);

 R_2 is H, C_1 - C_{12} alkyl (e.g., methyl), C_6 - C_{18} aryl (e.g., phenyl, napthaleneacetyl), C_1 - C_{12} acyl (e.g., formyl, acetyl, and myristoyl), C_7 - C_{18} aralkyl (e.g., benzyl), or C_7 - C_{18} alkaryl (e.g., p-methylphenyl);

 ${\rm A}^{25}$ is Arg, Lys, homo-Arg, diethyl-homo-Arg, Lys-NH-R (where R is H, a branched or straight chain ${\rm C}_1{\rm -C}_{10}$ alkyl group, or an aryl group), Orn, or is deleted;

 A^{26} is Ala, His, Thr, 3-Me-His, 1-Me-His, β -pyrozolylalanine, N-Me-His, Arg, Lys, homo-Arg, diethyl-homo-Arg, Lys- ϵ -NH-R (where R is H, a branched or straight chain C_1 - C_{10} alkyl group, or an aryl group), Orn, or is deleted;

A²⁷ is an aromatic amino acid;

A²⁸ is Leu, Ile, Val, Trp, Nle, Nva, Aib, Anb, or N-Me-Leu;

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A²⁹ is Asn, Ala, Gln, Gly, Trp, or N-Me-Asn; A³⁰ is Leu, Ile, Val, Trp, Nle, Nva, Aib, Anb, or N-Me-Leu;

A³¹ is Val, Ile, Trp, Nva, Aib, Anb, or N-Me-Val; A³² is Thr, Ser, N-Me-Ser, N-Me-Thr, or D-Trp;

 R_3 is H, C_1-C_{12} alkyl (e.g., methyl), C_6-C_{18} aryl (e.g., phenyl, napthaleneacetyl), C₁-C₁₂ acyl (e.g., formyl, acetyl, and myristoyl), C_7-C_{18} aralkyl (e.g., benzyl), or C_7 - C_{18} alkaryl

(e.g., p-methylphenyl); and

 R_4 is H, C_1 - C_{12} alkyl (e.g., methyl), C_6 - C_{18} aryl (e.g., phenyl, napthaleneacetyl), C_1-C_{12} acyl (e.g., formyl, acetyl, and myristoyl), C_7-C_{18} aralkyl (e.g., benzyl), or C7-C18 alkaryl (e.g., p-methylphenyl), or a pharmaceutically acceptable salt thereof.

In preferred embodiments A²⁷ is Phe, Nal, Bip, Pcp, Tic, Trp, Bth, Thi, or Dip.

In preferred embodiments Y is $A^{33}-A^{34}-A^{35}-A^{36}$

20 wherein

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A³³ is Arg, Lys, homo-Arg, diethyl-homo-Arg, Lys-NH-R (where R is H, a branched or straight ϵ – chain C_1-C_{10} alkyl group, or C_6-C_{18} aryl group), Cys, or Orn;

A³⁴ is Gln, Asn, Ala, Gly, N-Me-Gln, Aib, Cys, or Anb;

A³⁵ is Arg, Lys, homo-Arg, diethyl-homo-Arg, Lys-NH-R (where R is H, a branched or straight €chain C_1 - C_{10} alkyl group, or C_6 - C_{18} aryl. group), Cys, or Orn; and 30 -

> A³⁶ is an aromatic amino acid, Cys, or a pharmaceutically acceptable salt thereof.

Preferably, the compound has the formula $N-\alpha-Ac-$ Arg-His-Phe-Leu-Asn-Leu-Val-Thr-Arg-Gln-Arg-Tyr-NH2 (SEQ. 35 ID. NO. 8).

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In another aspect, the invention features novel dimeric analogs of peptide YY. The dimer may be formed by either including two peptides of Formula I, two peptides of Formula II, or one peptide of Formula I and 5 one peptide of Formula II. In one embodiment, the dimer is formed by utilizing a dicarboxylic acid linker capable of binding to a free amine, either primary or secondary, located within each peptide. See, e.g., R. Vavrek and J. Stewart, Peptides: Structure and Function 381-384 (Pierce 10 Chemical Co. 1983). Examples of suitable dicarboxylic acid linkers are succinic acid, glutamic acid, and phthalic acid. In other embodiments, the dimer is formed by utilizing an amino acid linker capable of binding to a free amine group of one peptide and a free carboxyl group 15 of the other peptide. Preferably, the amino acid linker is a non α -amino acid. Examples of suitable amino acid linkers are amino-caproic acid and amino-valeric acid. In yet another embodiment, the dimer is formed by a disulfide bridge between cysteines located within each 20 peptide. See, e.g., M. Berngtowicz and G. Piatsueda, Peptides: Structure and Function 233-244 (Pierce Chemical Co. 1985); F. Albericio, et al., Peptides 1990. 535 (ESCOM 1991).

In yet another aspect, the invention features

analogs of Formula I or Formula II having at least one
pseudopeptide bond between amino acid residues. By
"pseudopeptide bond" is meant that the carbon atom
participating in the bond between two residues is reduced
from a carbonyl carbon to a methylene carbon, i.e., CH2
NH; or less preferably that of CO-NH is replaced with any
of CH2-S, CH2-CH2, CH2-O, or CH2-CO. A pseudopeptide
peptide bond is symbolized herein by "Y". Preferably,
the pseudopeptide bonds are located between one or more
amino acid residues, e.g., A²⁸YA²⁹, A²⁹YA³⁰, A³⁰YA³¹,

A³¹YA³², A³²YA³³, A³³YA³⁴, A³⁴YA³⁵, or A³⁵YA³⁶. In addition,

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such pseudopeptide bond analogs can be used to form dimeric analogs as is described above. A detailed discussion of the chemistry of pseudopeptide bonds is given in Coy et al. (1988) Tetrahedron 44:835-841.

In yet another aspect, the invention features radiolabeled analogs of Formula I and Formula II.

Preferably, the analogs have a tyrosine iodinated on the phenyl ring at carbon position 3 or 5. The radioactive iodine is preferably I¹²⁵ or I¹²³. An example of the chemistry associated with iodinated tyrosine residues within peptides, see European Patent Application 0389180, herein incorporated by reference. Accordingly, radiolabeled PYY analogs can be used for imaging PYY receptors, e.g., for imaging cells containing PYY receptors.

The symbol X, Y, Z, A²², A²³, A²⁴, and the like; and Ser, Leu or the like, as found in a peptide sequence herein stands for an amino acid residue, i.e., =N-CH(R)-CO- when it is at the N-terminus, or -NH-CH(R)-CO-N= when it is at C-terminus, or -NH-CH(R)-CO- when it is not at the N- or C-terminus, where R denotes the side chain (or identifying group) of an amino acid or its residue. For example, R is -CH₂COOH for Asp, R is -H for Gly, R is -CH₂OH for Ser, R is -CH₃ for Ala and R is -CH₂CH₂CH₂CH₂NH₂ for Arg. Also, when the amino acid residue is optically active, it is the L-form configuration that is intended unless the D-form is expressly designated.

As set forth above and for convenience in describing this invention, the conventional and nonconventional abbreviations for the various amino acids are used. They are familiar to those skilled in the art; but for clarity are listed below. All peptide sequences mentioned herein are written according to the usual convention whereby the N-terminal amino acid is on the left and the C-terminal amino acid is on the right. A

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short line between two amino acid residues indicates a peptide bond.

Asp = D = Aspartic Acid

Ala = A = Alanine

5 Arg = R = Arginine

Asn = N = Asparagine

Cys = C = Cysteine

Gly = G = Glycine

Glu = E = Glutamic Acid

10 Gln = Q = Glutamine

His = H = Histidine

Ile = I = Isoleucine

Leu = L = Leucine

Lys = K = Lysine

15 Met = M = Methionine

Phe = F = Phenylalanine

Pro = P = Proline

Ser = S = Serine

Thr = T = Threonine

20 Trp = W = Tryptophan

Tyr = Y = Tyrosine

Val = V = Valine

Orn = Ornithine

Nal = 2-napthylalanine

25 Nva = Norvaline

Nle = Norleucine

Thi = 2-thienylalanine

Pcp = 4-chlorophenylalanine

Bth = 3-benzothienyalanine

30 Bip = 4,4'-biphenylalanine

Tic = tetrahydroisoquinoline-3-carboxylic acid

Aib = aminoisobutyric acid

Anb = α -aminonormalbutyric acid

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Dip = 2,2-diphenylalanine

Thz = 4-Thiazolylalanine

The compounds of the present invention can be provided in the form of pharmaceutically acceptable

5 salts. Examples of preferred salts are those with therapeutically acceptable organic acids, e.g., acetic, lactic, maleic, citric, malic, ascorbic, succinic, benzoic, salicylic, methanesulfonic, toluenesulfonic, trifluoroacetic, or pamoic acid, as well as polymeric acids such as tannic acid or carboxymethyl cellulose, and salts with inorganic acids, such as hydrohalic acids, e.g., hydrochloric acid, sulfuric acid, or phosphoric acid and the like.

In another aspect, the invention features one of
the above compounds and a pharmaceutically acceptable
carrier substance in a therapeutic composition capable of
decreasing excess intestinal water and electrolyte
secretion.

In preferred embodiments, the composition is in
the form of a liquid, pill, tablet, or capsule for oral
administration; a liquid capable of being administered
nasally as drops or spray or a liquid for intravenous,
subcutaneous, parenteral, intraperitoneal or rectal
administration. The therapeutic composition can also be
in the form of an oil emulsion or dispersion in
conjunction with a lipophilic salt such as pamoic acid,
or in the form of a biodegradable sustained-release
composition for subcutaneous or intramuscular
administration. For maximum efficacy, zero-order release
is desired.

In another aspect the invention features, a method for decreasing excess intestinal water and electrolyte secretion in a mammal, the method comprising administering to the mammal, e.g., a human, a

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therapeutically effective amount of the above mentioned compounds.

In addition, the invention features a method of regulating cell proliferation in a mammal, the method comprising administering to the mammal a therapeutically effective amount of the composition of the above mentioned compounds. Preferably, the method regulates the proliferation of an intestinal cell.

The invention also features methods for increasing nutrient transport, regulating lipolysis, and regulating blood flow in a mammal, the methods comprising administering to the mammal a therapeutically effective amount of the above mentioned compositions.

The compounds of the invention exhibit a broad

15 range of biological activities related to their
antisecretory and antimotility properties. The compounds
are believed to suppress gastrointestinal secretions by
direct interaction with epithelial cells or, perhaps, by
inhibiting secretion of hormones or neurotransmitters

20 which stimulate intestinal secretion. The compounds of
the invention may also control intestinal blood flow
which in turn may modulate intestinal hydrostatic
pressure in favor of net water absorption.

The compounds of the invention are especially
useful in the treatment of any number of gastrointestinal
disorders (see e.g., Harrison's Principles of Internal
Medicine, McGraw-Hill Inc., New York, 12th Ed.) that
are associated with excess intestinal electrolyte and
water secretion as well as decreased absorption, e.g.,
infectious (e.g., viral or bacterial) diarrhea,
inflammatory diarrhea, short bowel syndrome, or the
diarrhea which typically occurs following surgical
procedures, e.g., ileostomy. Examples of infectious
diarrhea include, without limitation, acute viral
diarrhea, acute bacterial diarrhea (e.g., salmonella,

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campylobacter, and clostridium or due to protozoal infections), or traveller's diarrhea (e.g., Norwalk virus or rotavirus). Examples of inflammatory diarrhea include, without limitation, malabsorption syndrome, 5 tropical spue, chronic pancreatitis, Crohn's disease, diarrhea, and irritable bowel syndrome. It has also been discovered that the peptides of the invention can be used to treat an emergency or life-threatening situation involving a gastrointestinal disorder, e.g., after 10 surgery or due to cholera. Furthermore, the compounds of the invention can be used to treat patients suffering from Acquired Immune Deficiency Syndrome (AIDS), especially during cachexia.

The compounds of the invention are also useful for 15 inhibiting small intestinal fluid and electrolyte secretion, augmenting nutrient transport -- as well as increasing cell proliferation -- in the gastrointestinal tract, regulating lipolysis in, e.g, adipose tissue, and regulating blood flow in a mammal.

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The compounds of the invention are advantageous because they are truncated versions of the natural PYY peptide; thus, the shorter peptide not only facilitates easier synthesis and purification of the compounds, but also improves and reduces manufacturing procedures and 25 expenses. Moreover, a shorter PYY compound is advantageous because such peptides will interact solely with PYY receptors and not with homologous receptors such as NPY Y1 and Y3; thus, minimizing unwanted agonist or antagonist side reactions.

Other features and advantages of the invention will be apparent from the following description of the preferred embodiments thereof, and from the claims.

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<u>Detailed Description</u>

The drawings will first be described.

Drawings

FIG. 1 shows a semipreparative reversed phase 5 chromatogram of N-α-Ac-[Phe²⁷]PYY(22-36) (SEQ. ID. NO. 3) (≈25mg) obtained by HF cleavage. Conditions: Vydac C18 semipreparative column (250 X 10mm, 300 Å pore size, 10 micron particle size); flow rate 4.7 ml/min; fractions 1, 2, 3, and 4 were collected and analyzed by analytical chromatography. The homogeneous fractions (1-3) were combined and dried in a speed vac.

FIG. 2 shows a graph of the inhibition of ¹²⁵IPYY binding to rat jejunal membranes by increasing
concentrations of PYY (SEQ. ID. NO. 1), PYY(22-36) (SEQ.

15 ID. NO. 10), [Im-DNP-His²⁶]PYY (SEQ. ID. NO. 9),
[Ala³²]PYY(22-36) (SEQ. ID. NO. 11), [Ala^{23,32}]PYY(22-36)
(SEQ. ID. NO. 12), [Glu²⁸]PYY(22-36) (SEQ. ID. NO. 13), Nα-Ac-PYY(22-36) (SEQ. ID. NO. 14), N-α-Ac-[p.ClPhe²⁸]PYY(22-36) (SEQ. ID. NO. 15), N-α-Ac-[Glu²⁶]PYY(2220 36) (SEQ. ID. NO. 16), N-α-Ac-[Phe²⁷]PYY(22-36) (SEQ. ID.
NO. 3), N-α-Ac-[N-Me-Tyr²⁶]PYY(22-36) (SEQ. ID. NO. 17), Nα-Myristoyl-PYY(22-36) (SEQ. ID. NO. 18), N-αNaphthaleneacetyl-PYY(22-36) (SEQ. ID. NO. 19), and PYY
(22-26) (SEQ. ID. NO. 10).

FIGS. 3A-B show the antisecretory effects of PYY (SEQ. ID. NO. 1), PYY(22-36) (SEQ. ID. NO. 10) and analogs up one baseline short circuit current (SCC) in voltage clamped preparation of rat jejunum. Values of changes in SCC are quoted of μ A/0.6cm², mean ±SEM from between 3 and 7 different jejunal preparations. Peptides shown in A and B are denoted by the same symbol as in FIG. 2.

FIG. 4 shows a graph of the inhibition of $^{125}I-PYY$ binding to rat jejunal membranes by increasing concentrations of PYY, $N-\alpha-Ac-PYY(22-36)$ (SEQ. ID. NO. 35 14), $N-\alpha-Ac-[Tic^{27}]PYY(22-36)$ (SEQ. ID. NO. 25), $N-\alpha-Ac-$

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[Bip²⁷]PYY(22-36) (SEQ. ID. NO. 22), N- α -Ac-[Nal²⁷]PYY(22-36) (SEQ. ID. NO. 23), N- α -Ac-[Bth²⁷]PYY(22-36) (SEQ. ID. NO. 21), N- α -Ac-[Phe²⁷]PYY(22-36) (SEQ. ID. NO. 3), N- α -Ac-[Phe²⁷]PYY(25-36) (SEQ. ID. NO. 26), N- α -Ac-[Trp²⁷]PYY(22-56) (SEQ. ID. NO. 5), and N- α -Ac-[Thi²⁷]PYY(22-36) (SEQ. ID. NO. 6).

There now follows a description of the synthesis, analysis for biological efficacy and use of the preferred embodiments of the invention. In order to determine the structural requirements necessary to elicit antisecretory effects, several analogs of the PYY active site, PYY(22-36), were synthesized and their binding and antisecretory potencies in rat jejunum were compared.

We now describe the structure, synthesis, and use of preferred embodiments of the invention.

STRUCTURE

The peptides of the invention have the general formula recited in the Summary of the Invention above.

They all have an aromatic amino acid group at position 27 which is important for both antisecretory activity and utility as antidiarrheal compounds.

SYNTHESIS

The peptides of the present invention may be synthesized by any techniques that are known to those skilled in the peptide art. An excellent summary of the many techniques so available may be found in *Solid Phase Peptide Synthesis* 2nd ed. (Stewart, J.M. and Young, J. D. Pierce Chemical Company, Rockford, IL, 1984).

The peptides listed in Table 1 and Table 2 were synthesized as follows. Peptide synthesis was performed on an Applied Biosystems Model 430A synthesizer. Amino acid and sequence analyses were carried out using Waters Pico-Tag and Applied Biosystems Model 470A instruments,

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respectively. Peptides were purified using a Waters
Model 600 solvent delivery system equipped with a Model
481 Spectrophotometer and U6K injector according to
standard protocols. Peptide masses were determined at
the University of Michigan, Protein Chemistry Facility,
Ann Arbor, Michigan according to standard methods. All
Boc-L-amino acid derivatives, solvents, chemicals and the
resins were obtained commercially and used without
further purification.

Paramethylbenzhydroxylamine (MBHA) resin (0.45 10 mmol, -NH2) was placed in the reaction vessel of the peptide synthesizer and the protected amino acid derivatives were sequentially coupled using the program provided by the manufacturers modified to incorporate a 15 double coupling procedure (see, e.g., Balasubramaniam et al., Peptide Research 1: 32, 1988). All amino acids were coupled using 2.2 equivalents of preformed symmetrical anhydrides. Arg, Gln and Asn, however, were coupled as preformed 1-hydroxybenzotriazole (HOBT) esters to avoid 20 side reactions. At the end of the synthesis, the N- α -Boc group was removed and in some instances the free $\alpha-NH2$ was acetylated by reaction with acetic anhydride (2 equivalents) and diisopropyl ethylamine until a negative ninhydrin test was obtained (Anal. Biochem. 34:595, 25 1970). The peptide resin (~1.0 g) was then treated with HF (10 ml) containing p-cresol (~0.8 g) for 1 h at -2 to -4 °C. The HF was evacuated and the residue was transferred to a fritted filter funnel with diethyl ether, washed repeatedly with diethyl ether, extracted 30 with acetic acid (2 X 15 ml) and lyophilized. peptides thus obtained were purified by semipreparative RP-HPLC as shown in Fig. 1.

Examples of the synthesized analogs are:

[im-DNP-His²⁶]PYY
35 YPAKPEAPGEDASPEELSRYYASLR [im-DNP-His²⁶]YLNLVTRQRY-NH₂ (SEQ. ID No. 9)
PYY(22-36)

 $\mathbb{R}_{2} + \mathbb{R}_{2} + \mathbb{R}_{3} + \mathbb{R}_{3} + \mathbb{R}_{3} = \mathbb{R}_{3}$

	ASLRHYLNLVTRQRY-NH ₂	(SEQ. ID No. 10)
	[Ala ³²]PYY ASLRHYLNLV [Ala] RQRY-NH ₂	(SEQ. ID No. 11)
5	$(Ala^{23,32})PYY$ A (Ala) L R H Y L N L V (Ala) R Q R Y-NH ₂	(SEQ. ID No. 12)
	[Glu ²⁸]PYY(22-36) A S L R H Y [Glu] N L V T R Q R Y-NH ₂	(SEQ. ID No. 13)
	N-α-Ac-PYY(22-36) N-α-Ac-A S L R H Y L N L V T R Q R Y-NH ₂	(SEQ. ID No. 14)
10	N-α-Ac[p.Cl.Phe ²⁶]PYY N-α-Ac-A S L R [p.Cl.Phe ²⁶] Y L N L V T R Q R Y-NH ₂	(SEQ. ID No. 15)
	N-α-Ac[Glu ²⁸]PYY N-α-Ac-A S L R H Y [Glu] N L V T R Q R Y-NH ₂	(SEQ. ID No. 16)
15	N-a-Ac [Phe ²⁷] PYY	_
15	N-α-Ac-A S L R H [Phe] E N L V T R Q R [N-Me-Tyr]-NH ₂ N-α-Ac[N-Me-Tyr ³⁶]PYY	(SEQ. ID No. 3)
	N-α-Ac-A S L R H Y E N L V T R Q R [N-Me-Tyr]-NH ₂ N-α-myristoyl-PYY(22-36)	(SEQ. ID No. 17)
20	N-α-myristoyl-A S L R H Y L N L V T R Q R Y-NH ₂ N-α-naphthaleneacetyl-PYY(22-36)	(SEQ. ID No. 18)
	N-α-naphthaleneacetyl -A S L R H Y L N L V T R Q R Y-NH ₂ N-α-Ac[Phe ²⁷] PYY	(SEQ. ID No. 19)
	N-α-Ac-A S L R H [Phe] E N L V T R Q R (N-Me-Tyr)-NH ₂ N-α-Ac-PYY(22-36)	(SEQ. 1D No. 3)
25	N-α-AC-A S L R H Y L N L V T R Q R Y-NH ₂ N-α-AC-[Bth ²⁷]PYY(22-36)	(SEQ. ID No. 20)
	N-α-AC-A S L R H [Bth] L N L V T R Q R Y-NH ₂	(SEQ. ID No. 21)
20	N-α-AC-A S L R H [Bip] L N L V T R Q R Y-NH ₂ N-α-AC-[Nal ²⁷]PYY(22-36)	(SEQ. 1D No. 22)
30	N-α-AC-A S L R H [Nal] L N L V T R Q R Y-NH2	(SEQ. ID No. 23)
	N-α-Ac-[Trp ²⁷]PYY(22-36) N-α-Ac-A S L R H [Trp] L N L V T R Q R Y-NH ₂	(SEQ. ID No. 5)
35	N-α-Ac-[Thi ²⁷]PYY(22-36) N-α-Ac-A S L R H [Thi] L N L V T R Q R Y-NH ₂	(SEQ. ID No. 6)
	N-α-Ac-[Tic ²⁷]PYY(22-36) N-α-Ac-A S L R H [Tic] L N L V T R Q R Y-NH ₂	(SEQ. ID No. 25)
	N-α-Ac-[Phe ²⁷]PYY(25-36) N-α-Ac-H [Phe] L N L V T R Q R Y-NH ₂	(SEQ. ID No. 26)
40	N-α-Ac-[Phe ²⁷ , Thi ³⁶] PYY(22-36) N-α-Ac-A S L R H [Phe] L N L V T R Q R [Thi]-NH ₂	(SEQ. ID No. 27)
	N-α-Ac-[Thz ²⁶ ,Phe ²⁷]PYY(22-36) N-α-Ac-A S L R [Thz] [Phe] L N L V T R Q R Y-NH ₂	(SEQ. ID No. 28)
45	N-α-Ac-[Pcp ²⁷]PYY(22-36) N-α-Ac-A S L R H [Pcp] L N L V T R Q R Y-NH ₂	(SEQ. ID No. 29)
	N-α-Ac-[Phe ^{22,27}]PYY(22-36) N-α-Ac-[Phe] S L R H [Phe] L N L V T R Q R Y-NH ₂	(SEQ. ID No. 30)

	N-α-Ac-[Tyr ²² ,Phe ²⁷]PYY(22-36) N-α-Ac-[Tyr] S L R H [Phe] L N L V T R Q R Y-NH ₂	(SEQ.	ID No.	. 7)
	N-α-Ac-[Trp ²⁸]PYY(22-36) N-α-Ac- A S L R H Y [Trp] N L V T R Q R Y-NH ₂	(SEQ.	ID No.	. 31)
5	N-α-Ac-[Trp ³⁰]PYY(22-36) N-α-Ac- A S L R H Y L N [Trp] V T R Q R Y-NH ₂	(SEQ.	ID No.	. 32)
	N- α -Ac-[Ala 26 ,Phe 27]PYY(22-36) N- α -Ac- A S L R [Ala] [Phe] L N L V T R Q R Y-NH $_2$	(SEQ.	ID No.	. 33)
10	N-α-Ac-[Bth ²⁷]PYY(22-36) N-α-Ac- A S L R H [Bth]	(SEQ.	ID No.	. 34)
	N-α-Ac-[Phe ²⁷]PYY(22-36) N-α-Ac- A S L R H [Phe] L N L V T R Q R Y-NH ₂	(SEQ.	ID No.	. 35)
	N-α-Ac-[Phe ^{27,36}]PYY(22-36) N-α-Ac- A S L R H [Phe] L N L V T R Q R [Phe]-NH ₂	(SEQ.	ID No.	36)
15	N-α-Ac-[Phe ²⁷ , D-Trp ³²]PYY(22-36) N-α-Ac- A S L R H [Phe] L N L V [D-Trp] R Q R Y-NH ₂	(SEQ.	ID No.	. 37)

ANALYSIS

Binding Studies

Preparation of ¹²⁵I-PYY labeled only at Tyr³⁶ and rat jejunal epithelial plasma membranes were performed according to standard methods (see, e.g., Laburthe et al. Endocrinology, supra; Servin et al. supra; Voisin et al. Ann. N. Y. Acad. Sci. 611:343, 1990). Binding experiments were conducted in a total volume of 0.25 ml 60 mM HEPES buffer, pH 7, containing 2% BSA, 0.1% bacitracin, 5 mM MgCl₂ and 0.05 nM ¹²⁵I-PYY with or without competing peptides. Bound and free peptides were separated by centrifugation at 20,000 X g for 10 min. Non-specific ¹²⁵I-PYY binding was determined in the presence of 1 μM unlabeled PYY represented 10% of the total binding.

Short Circuit Current Measurements

The antisecretory effects of the peptides were investigated by measuring the short-circuit current (SCC) in rat jejunal mucosa mounted in a Ussing chamber and automatically voltage clamped as described by Cox et al. (J. Physiol. supra). Briefly, strips of mucosa were placed between two halves of perspex Ussing chambers

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(window size, 0.6 cm²) containing oxygenated (95% O₂/5% CO₂) Krebs-Henseleit solution (NaCl, 117 mM, KCl 4.7 mM, CaCl₂, 2.5 mM; MgSO₄ 1.2 mM, NaHCO₃ 24.8 mM and glucose 11.1 mM), pH 7.4, 37°C. Routinely, four preparations of jejunum were obtained from each animal and these exhibited comparable potential differences and SCC, but they were not paired. Preparations were automatically voltage clamped using a W-P dual voltage clamp and the SCC displayed continuously on pen recorders. Once a stable baseline SCC was reached, peptides were added to the basolateral reservoir only, and cumulative concentration-response profiles constructed.

Data Analyses

All points in the binding experiments are the mean of at least three experiments performed in duplicate. For clarity, the SEMs in the binding experiments are not shown in Fig. 2, but were less than 10%. Values of changes in SCC are quoted as $\mu A/0.6 \text{cm}^2$ mean \pm 1 SEM from between 3 and 7 different preparations. EC_{50} values were calculated from pooled cumulative concentration - response curves using an iterative curve fitting program. Comparison of data groups (SCC recordings) were made using unpaired Student's t-tests where a p value <0.5 was considered statistically significant.

There now follows the results of the biological activities of the compounds of the invention (see Table 1 and Table 2). As described below, the tested compounds were assayed for purity and for their binding and antisecretory potencies in rat jejunum.

Purified peptides were found to be > 96% homogeneous by analytical reversed phase chromatography and, in addition, had the expected amino acid composition and masses. For example, Fig. 1 shows the RP-HPLC chromatogram of N-α-Ac-[Phe²⁷]PYY(22-36)(SEQ. ID. NO. 3).

The free peptides were further characterized by sequence

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analysis (see, Table 1 and Table 2). The overall yields of the peptides were in the range of 10% to 30%.

PYY, [im-DNP-His²⁶]PYY (SEQ. ID. NO. 9) and the analogs of PYY(22-36)(SEQ. ID. NO. 10) displaced 125I-PYY 5 bound to rat jejunal epithelial plasma membranes in a concentration-dependent manner. Although [im-DNP- His^{26} PYY (SEQ. ID. NO. 9) and PYY(22-36) (SEQ. ID. NO. 10) were 20-times less potent than PYY based on IC50 values, they displayed the same maximal response as the 10 intact hormone (Fig. 2, Table 1). Substitution of Thr 32 with Ala as in [Ala³²]PYY(22-36)(SEQ. ID. NO. 11) resulted in the lowering of the binding potency while the replacement of both Ser²³ and Thr³² with Ala further reduced the receptor affinity. Also, introduction of a 15 negative charge at position 28 without altering the helicity as in $[Glu^{28}]PYY(22-36)$ (SEQ. ID. NO. 13) decreased the binding possibly due to the disruption of the ionic interactions. Since the hydrophobic groups are known to increase the interaction with the receptors 20 (Balasubramaniam et al. Biochem. Biophys. Res. Comm. 137:1041, 1986), the binding of a N- α -myristoyl- and N- α naphthaleneacetyl-derivatives of PYY(22-36) was also determined. Both these analogs exhibited slightly lower binding affinity than PYY(22-36)(SEQ. ID. NO. 10) 25 possibly due to increased steric hinderance. On the other hand, $N-\alpha$ -acetylation of PYY(22-36) (SEQ. ID. NO. 14) increased the receptor affinity four times. Further structure-activity studies with $N-\alpha-Ac-PYY(22-36)$ (SEQ. ID. NO. 20) revealed that substitution of Tyr36 with N-Me-30 Tyr or His^{26} with p.Cl-Phe lowers the binding potency. However, replacement of Tyr27 with Phe increased the receptor affinity by 28%. As a control, the binding of PYY(22-36)(SEQ. ID. NO. 10) and several of its analogs were also tested. However, none of these analogs 35 inhibited the binding of ^{125}I -PYY even at 10 μ M.

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In mucosal preparations of rat jejunum PYY(22-36) (SEQ. ID. NO. 10) analogs reduced the baseline SCC in a concentration dependent manner (Fig. 3A and B) and calculated EC₅₀ values are listed in Table 1. The PYY(22-5 36) (SEQ. ID. NO. 10) analogs were generally less potent as antisecretory agents than as inhibitors of binding. The order of analog potency was similar to that from binding studies with two notable exceptions, namely N- α myristoyl-PYY(22-36) (SEQ. ID. NO. 18) and N- α -10 naphthaleneacetyl-PYY(22-36) (SEQ. ID. NO. 19). acetylation and substitution of Tyr²⁷ with Phe increased the antisecretory potency of PYY(22-36) and this analog, $N-\alpha-Ac-[Phe^{27}]$ PYY(22-36) (SEQ. ID. NO. 3), was only 9times less potent than the intact hormone. Furthermore, 15 there was no significant difference between the maximal inhibitory responses, these being - 12.6±2.4 and - $12.0\pm1.3\mu$ A/0.6cm² for PYY (440 nM, n = 6) (SEQ. ID. NO. 1) and $N-\alpha-Ac-[Phe^{27}]$ PYY(22-36) (1.4 μ M, n = 7) (SEQ. ID. NO. 3), respectively.

TABLE 1: Comparison of the binding and antisecretory potencies of PYY, PYY fragments and their analogs

5	PEPTIDES RT ^a MH+ (Calc.)	BINDING ^b (min)	scc ^b	IC ₅₀ (nM)	EC ₅₀ (nM)
	PYY (SEQ. ID. NO. 1)	4.8	4240.2 (4241.7)	0.2	1.7
	NPY (SEQ. ID. NO. 24)	34.0 ^c	4253.8 (4254.7)	2.0	9 ^d
	[im-DNP-His ²⁶]PYY (SEQ. ID. NO. 9)	8.7 ^c	4406.9 (4407.8)	4.0	72
10	PYY(22-36) (SEQ. ID. NO. 10)	4.4	1888.8 (1890.2)	4.0	77
	[Ala ³²]PYY(22-36) (SEQ. ID. NO. 11)	4.7	1858.8 (1860.2)	71	n.d.
	[Ala ^{23,32}]PYY(22-36) (SEQ. ID. NO. 12)	4.3	1842.8 (1844.2)	>10,000	n.d.
	[Glu ²⁸]PYY(22-36) (SEQ. ID. NO. 13)	3.8	1905.1 (1906.2)	199	n.d.
	N-α-Ac-PYY(22-36) (SEQ. ID. NO. 14)	10.0	1930.9 (1932.2)	1.12	40
15	N-α-Ac-[p.ClPhe ²⁶]PYY(22-36) (SEQ. ID. NO. 15)	14.9 ^c	1975.4 (1976.7)	50	124
	N-α-Ac-[Glu ²⁸]PYY(22- 3 6) (SEQ. ID. NO. 16)	3.9	1947.0 (1948.2)	44.7	3,000
20	N-α-Ac-[N-Me-Tyr ³⁶]PYY(22-36) (SEQ. ID. NO. 17)	13.5	1945.3 (1946.3)	354	792
<i>:</i> ,	N-α-Ac-[Phe ²⁷]PYY(22-36) (SEQ. ID. NO. 3)	8.3	1915.3 (1916.2)	0.80	15.1
•. •			e 2.74 1.74		
	. N-α-Myristoyl-PYY(22-36) (SEQ. ID. NO. 18)	4.8	2099.0 (2100.6)	17.8	3,300
25	N-α-Naphthaleneacetyl-PYY(22-36) (SEQ. ID. NO. 19)	17.0	2056.9 (2058.4)	8.9	19,500

a: isocratic, 27% CH₃CN containing 0.1% TFA; b: mean of three separate experiments; c: isocratic, 32% CH₃CN containing 0.1% TFA; d: from reference 10; n.d.: not determined

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 $N-\alpha$ -myristoyl-PYY(22-36)(SEQ. ID. NO. 18) and $N-\alpha$ naphthaleneacetyl-PYY(22-36) (SEQ. ID. NO. 19) analogs, in contrast to their moderate binding potency, exhibited poor antisecretory responses with threshold 5 concentrations of about 20nM and EC₅₀ values greater than 2 and 30 μ M respectively. After a cumulative concentration of 7.4 μ M, N- α -myristoyl-PYY(22-36) (SEQ. ID. No. 18) reduced the basal SCC by - $5.2\pm0.6\mu\text{A}/0.6\text{cm}^2$ (n = 7). Subsequent addition of PYY (100 nM) further 10 reduced the SCC by $-10.2\pm0.7\mu\text{A}/0.6\text{cm}^2$ (n = 7) and this was not significantly different from control responses to PYY(22-36) (SEQ. ID. 10) could antagonize PYY responses, three tissues were treated with the analog (1 μ M) and PYY concentration-15 response curves were constructed and compared with controls. The fragment reduced the basal current by - $0.4\pm0.3~\mu\text{A}/0.6\text{cm}^2$ and the resultant PYY EC₅₀ value $(4.4\pm1.2 \text{ nM}, \text{n} = 3)$ did not differ significantly from that of the nontreated controls $(2.6\pm1.1 \text{ nM}, n = 3)$. These results show that modification of the active 20 site of PYY (SEQ. ID. NO. 1), PYY(22-36)(SEQ. ID. NO. 10), can lead to a substantial increase in both the binding and antisecretory potencies of this fragment. The key analogs in this series exhibited the following 25 order of potency: PYY (SEQ. ID. NO. 1) > N- α -Ac- $[Phe^{27}]PYY(22-36)$ (SEQ. ID. NO. 3) > N-\alpha-Ac-PYY(22-36) (SEQ. ID. NO. 14) > PYY(22-36) (SEQ. ID. NO. 10). Furthermore, our investigations revealed that the hydroxyl groups of Ser^{23} and Thr^{32} as well as the imidazole group of His^{26} 30 are important for interaction with intestinal PYYpreferring receptors. Although there was, in general, a good correlation between the binding and antisecretory potencies of the analogs, there were also notable exceptions.

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N-α-myristoyl-PYY(22-36) (SEQ. ID. NO. 18) and N-α-naphthaleneacetyl-PYY(22-36) (SEQ. ID. NO. 19) analogs inhibited ¹²⁵I-PYY binding with moderate potency, but exhibited poor antisecretory responses. This observation suggested that these analogs may be antagonists. However, prior pretreatment of jejunal membranes with these analogs failed to significantly alter the antisecretory responses to PYY and the reason for the discrepancy remains unclear at present.

Table 2 and Fig. 4 present the IC_{50} values for additional PYY(22-36) (SEQ. ID. NO. 10) and PYY (25-36) analogs. Based on the results presented in Table 2 the analogs in this series exhibited the following order of potency:

15 $N-\alpha-Ac-[Tic^{27}]PYY(22-36)$ (SEQ. ID. NO. 25) $< N-\alpha-Ac-[Bip^{27}]PYY(22-36)$ (SEQ. ID. NO. 22) $< N-\alpha-Ac-[Nal^{27}]PYY(22-36)$ (SEQ. ID. NO. 23) $< N-\alpha-Ac-[Bth^{27}]PYY(22-36)$ (SEQ. ID. NO. 21) $< N-\alpha-Ac-[Phe^{27}]PYY(22-36)$ (SEQ. ID. NO. 3) $< N-\alpha-Ac-[Phe^{27}]PYY(25-20)$ 36) (SEQ. ID. NO. 26) $< N-\alpha-Ac-[Trp^{27}]PYY(22-36)$ (SEQ. ID. NO. 5) $< N-\alpha-Ac-[Thi^{27}]PYY(22-36)$ (SEQ. ID. NO. 6) $< N-\alpha-Ac-PYY(22-36)$ (SEQ. ID. NO. 14) < PYY (SEQ. ID. NO. 1).

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Comparison of Receptor Binding Data for PYY and PYY analogs

	PEPTIDE NO.	Peptide Structure	IC ₅₀ (nM)		
		PYY (SEQ. ID. NO. 1)	0.04		
i		$N-\alpha-Ac-PYY(22-36)$ (SEQ. ID. NO. 14)	0.08		
	905	$N-\alpha-Ac-[Bth^{27}]PYY(22-36)$ (SEQ. ID. NO. 21)	0.22		
5	906	$N-\alpha-Ac-[Bip^{27}]PYY(22-36)$ (SEQ. ID. NO. 22)	4.46		
	911	$N-\alpha-Ac-[Nal^{27}]PYY(22-36)$ (SEQ. ID. NO. 23)	0.39		
	915	N-α-Ac-[Trp ²⁷]PYY(22-36) (SEQ. ID. NO. 5)	0.10		
	916	$N-\alpha-Ac-[Thi^{27}]PYY(22-36)$ (SEQ. ID. NO. 6)	0.095		
	914	N-α-Ac-[Phe ²⁷]PYY(25-36) (SEQ. ID. NO. 26)	0.15		
10	913	$N-\alpha-Ac-[Tic^{27}]PYY(22-36)$ (SEQ. ID. NO. 25)	4.50		

10

NPY/PYY receptors characterized to date have been broadly classified into Y-1, Y-2 and Y-3 subtypes (Balsubramaniam et al. J. Biol. Chem. 265:14724, 1990; Michel, Trends Pharmacol. Sci. 12:389, 1991). Both Y-1 15 and Y-2 receptors exhibit a preference for PYY over NPY, and more significantly C-terminal fragments of NPY and PYY are effective only at the Y-2 subtype. Y-3 receptors, on the other hand, exhibit a greater affinity for NPY than PYY. Since rat jejunal mucosa antisecretory 20 responses show an order of agonist potency PYY (SEQ. ID. NO. 1) > NPY (SEQ. ID. NO. 24) > PYY(13-36)(SEQ. ID. NO. 32) > NPY(13-36)(SEQ. ID. NO. 33) these epithelial receptors are Y-2 like, and are completely insensitive to the Y-1 selective agonist [Pro34]NPY (Cox et al. Peptides, 25 supra). The results further describe $N-\alpha-Ac-PYY(22-36)$

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(SEQ. ID. NO. 14) and N- α -Ac-[Phe²⁷]PYY(22-36)(SEQ. ID. NO. 3) to be more potent than PYY(22-36)(SEQ. ID. NO. 10) and the corresponding C-terminal fragments of NPY of varying lengths (Cox et al. Br. J. Pharmacol. supra).

- 5 The higher affinity for PYY (SEQ. ID. NO. 1) and its Cterminal fragments compared with NPY (SEQ. ID. NO. 24) and its respective fragments is in agreement with the order of potency obtained from receptor binding studies with rat intestinal epithelial membranes (Laburthe et al.
- 10 supra; Laburthe, supra; Voisin et al. Ann. N.Y. Acad. Sci. supra; Voisin et al. Am. J. Physiol.)

In addition, analogs listed in Table 3 were synthesized as described above and tested for binding activity. The results shown in Table 3 demonstrate that $N-\alpha-Ac-[Tyr^{22}, Phe^{27}]PYY(22-36)$ (SEQ. ID. NO. 7) is similar in its competitive binding as PYY (SEQ. ID. NO. 1), indicating that the introduction of an aromatic amino acid, e.g., Tyr, at position 22 is an effective PYY analog.

20. TABLE 3

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PEPTIDE NO.	EPTIDE NO. Peptide Structure	
	PYY (SEQ. ID. NO. 1)	0.10
917	N-α-Ac-[Phe ²⁷ , Thi ³⁶]PYY(22-26) (SEQ. ID. NO. 27)	4.46
918	$N-\alpha-Ac-[Thz^{26}, Phe^{27}]Pyy(22-36)$ (SEQ. ID. NO. 28)	4.50
904	N-α-Ac-[Pcp ²⁷]PYY(22-36) (SEQ. ID. NO. 29)	1.58
908	N-α-Ac-[Phe ^{22,27}]PYY(22-36) (SEQ. ID. NO. 30)	11.22
910	$N-\alpha-Ac-[Tyr^{22}, Phe^{27}]PYY(22-36)$ (SEQ. ID. NO. 7)	0.10

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<u>USE</u>

In the practice of the method of the present invention, an effective amount of an any one or combination of the analogs of the invention, e.g., $N-\alpha$ -5 Ac-[Phe²⁷]PYY(22-36)(SEQ. ID. NO. 3), N- α -Ac- $[Trp^{27}]PYY(22-36)$ (SEQ. ID. NO. 24), $N-\alpha-Ac-[Phe^{27}]PYY(25-$ 36) (SEQ. ID. NO. 3), $N-\alpha-Ac-[Thi^{27}]PYY(22-36)$ (SEQ. ID. NO. 6) or derivative thereof, is administered via any of the usual and acceptable methods known in the art, either 10 singly or in combination with another compound or compounds of the present invention. These compounds or compositions can thus be administered orally (e.g., buccal cavity), sublingually, parenterally (e.g., intramuscularly, intravenously, or subcutaneously), 15 rectally (e.g., by suppositories or washings), transdermally (e.g., skin electroporation) or by inhalation (e.g., by aerosol), and in the form or either solid, liquid or gaseous dosage, including tablets and suspensions. The administration can be conducted in a 20 single unit dosage form with continuous therapy or in a single dose therapy ad libitum.

Thus, the method of the present invention is practiced when relief of symptoms is specifically required or perhaps imminent. Alternatively, the method of the present invention is effectively practiced as continuous or prophylactic treatment.

Useful pharmaceutical carriers for the preparation of the compositions hereof, can be solids, liquids or gases; thus, the compositions can take the form of tablets, pills, capsules, suppositories, powders, enterically coated or other protected formulations (e.g. binding on ion-exchange resins or packaging in lipid-protein vesicles), sustained release formulations, solutions, suspensions, elixirs, aerosols, and the like.

The carrier can be selected from the various oils

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including those of petroleum, animal, vegetable or synthetic origin, e.g., peanut oil, soybean oil, mineral oil, sesame oil, and the like. Water, saline, aqueous dextrose, and glycols are preferred liquid carriers, 5 particularly (when isotonic with the blood) for injectable solutions. For example, formulation for intravenous administration comprise sterile aqueous solutions of the active ingredient(s) which are prepared by dissolving solid active ingredient(s) in water to 10 produce an aqueous solution, and rendering the solution sterile. Suitable pharmaceutical excipients include starch, cellulose, talc, glucose, lactose, talc, gelatin, malt, rice, flour, chalk, silica, magnesium stearate, sodium stearate, glycerol monostearate, sodium chloride, 15 dried skim milk, glycerol, propylene glycol, water, ethanol, and the like. The compositions may be subjected to conventional pharmaceutical additives such as preservatives, stabilizing agents, wetting or emulsifying agents, salts for adjusting osmotic pressure, buffers and 20 the like. Suitable pharmaceutical carriers and their formulation are described in Remington's Pharmaceutical Sciences by E.W. Martin. Such compositions will, in any event, contain an effective amount of the active compound together with a suitable carrier so as to prepare the 25 proper dosage form for proper administration to the recipient.

The dose of the compound of the present invention for treating the above-mentioned disorders varies depending upon the manner of administration, the age and the body weight of the subject, and the condition of the subject to be treated, and ultimately will be decided by the attending physician or veterinarian. Such amount of the active compound as determined by the attending physician or veterinarian is referred to herein as a "therapeutically effective amount". Thus, a typical

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administration is oral administration or parenteral administration. The daily dose in the case of oral administration is typically in the range of 0.1 to 100 mg/kg body weight, and the daily dose in the case of parenteral administration is typically in the range of 0.001 to 50 mg/kg body weight.

To be effective for the prevention or treatment of gastroenterological disorders, especially infectious (e.g. viral or bacterial) or inflammatory diarrhea, or diarrhea resulting from surgery, it is important that the therapeutic agents be relatively non-toxic, non-antigenic and non-irritating at the levels in actual use.

It is understood that the examples and embodiments described herein are for illustrative purposes only and that various modifications or changes in light thereof will be suggested to persons skilled in the art and are to be included within the spirit and purview of this application and scope of the appended claims.

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SEQUENCE LISTING

- (1) GENERAL INFORMATION
- (i) APPLICANT: UNIVERSITY OF CINCINNATI
- (ii) TITLE OF THE INVENTION: ANALOGS OF PEPTIDE YY AND USES THEREOF
 - (iii) NUMBER OF SEQUENCES: 30
 - (iv) CORRESPONDENCE ADDRESS:
 - (A) ADDRESSEE: Fish & Richardson P.C.
 - (B) STREET: 225 Franklin Street (C) CITY: Boston

 - (D) STATE: MA
 - (E) COUNTRY: US
 - (F) ZIP: 02110-2804
 - (v) COMPUTER READABLE FORM:

 - (A) MEDIUM TYPE: Diskette
 (B) COMPUTER: IBM Compatible
 - (C) OPERATING SYSTEM: DOS
 - (D) SOFTWARE: FastSEQ Version 2.0
 - (vi) CURRENT APPLICATION DATA:
 - (A) APPLICATION NUMBER:
 - (B) FILING DATE: 13-NOV-1996 (C) CLASSIFICATION:
 - (viii) ATTORNEY/AGENT INFORMATION:
 - (A) NAME: Tsao, Y. Rocky
 - (B) REGISTRATION NUMBER: 34,476
 - (C) REFERENCE/DOCKET NUMBER: 00537/105W01
 - (ix) TELECOMMUNICATION INFORMATION:
 - (A) TELEPHONE: 617-542-8006
 - (B) TELEFAX: 617-542-8906 (C) TELEX:

 - (2) INFORMATION FOR SEQ ID NO:1:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 36 amino acids (B) TYPE: amino acid

 - (C) STRANDEDNESS: N/A
 - (D) TOPOLOGY: linear
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

Tyr Pro Ala Lys Pro Glu Ala Pro Gly Glu Asp Ala Ser Pro Glu Glu 10 Leu Ser Arg Tyr Tyr Ala Ser Leu Arg His Tyr Leu Asn Leu Val Thr 20 Arg Gln Arg Tyr 35

- (2) INFORMATION FOR SEQ ID NO:2:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 36 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: N/A

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- (D) TOPOLOGY: linear
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:

Tyr Pro Ile Lys Pro Glu Ala Pro Gly Glu Asp Ala Ser Pro Glu Glu 10 Leu Asn Arg Tyr Tyr Ala Ser Leu Arg His Tyr Leu Asn Leu Val Thr Arg Gln Arg Tyr

- (2) INFORMATION FOR SEQ ID NO: 3:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 15 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: N/A
 - (D) TOPOLOGY: linear
- (ix) FEATURE:

(D) OTHER INFORMATION: Xaa in position 15 is an abbreviation of N-Me-Tyr. The sequence has an acetylated N-terminus (i.e., N- α -Ac), rather than an amino N-terminus (i.e., H2N-). The sequence has an amide C-terminus (i.e., CO-NH2), rather than a carboxyl C-terminus (i.e., CO-OH).

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:

Ala Ser Leu Arg His Phe Leu Asn Leu Val Thr Arg Gln Arg Xaa

- (i) SEQUENCE CHARACTERISTICS: (2) INFORMATION FOR SEQ ID NO:4:
 - - (A) LENGTH: 15 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: N/A (D) TOPOLOGY: linear
 - (ix) FEATURE:

(D) OTHER INFORMATION: The sequence has an amide Cterminus (i.e., CO-NH2), rather than a carboxyl C-terminus (i.e., CO-OH).

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:

Ala Ser Leu Arg His Phe Leu Asn Leu Val Thr Arg Gln Arg Tyr 1 5 10

- (2) INFORMATION FOR SEQ ID NO:5:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 15 amino acids (B) TYPE: amino acid

 - (C) STRANDEDNESS: single

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- (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: None
- (ix) FEATURE:

(D) OTHER INFORMATION: The sequence has an acetylated Nterminus (i.e., $N-\alpha-Ac$), rather than an amino N-terminus (i.e., H2N-). The sequence has an amide C-terminus (i.e., CO-NH2), rather than a carboxyl C-terminus (i.e., CO-OH).

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:

Ala Ser Leu Arg His Trp Leu Asn Leu Val Thr Arg Gln Arg Tyr

- (2) INFORMATION FOR SEQ ID NO:6:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 15 amino acids (B) TYPE: amino acid

 - (C) STRANDEDNESS: N/A
 - (D) TOPOLOGY: linear
- (ix) FEATURE:

(D) OTHER INFORMATION: Xaa in position 6 is an abbreviation of Thi (2-thienylalanine). The sequence has an acetylated N-terminus (i.e., N- α -Ac), rather than an amino N-terminus (i.e., H2N-). The sequence has an amide C-terminus (i.e., CO-NH2), rather than a carboxyl C-terminus (i.e., CO-OH).

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:

Ala Ser Leu Arg His Xaa Leu Asn Leu Val Thr Arg Gln Arg Tyr 15

- (2) INFORMATION FOR SEQ ID NO:7:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 15 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: N/A
 - (D) TOPOLOGY: linear
- (ix) FEATURE:

(D) OTHER INFORMATION: The sequence has an acetylated Nterminus (i.e., $N-\alpha-Ac$), rather than an amino N-terminus (i.e., H2N-). The sequence has an amide C-terminus (i.e., CO-NH2), rather than a carboxyl C-terminus (i.e., CO-OH).

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:

Tyr Ser Leu Arg His Phe Leu Asn Leu Val Thr Arg Gln Arg Tyr

- (2) INFORMATION FOR SEQ ID NO:8:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 12 amino acids

- (B) TYPE: amino acid
- (C) STRANDEDNESS: N/A
- (D) TOPOLOGY: linear

(ix) FEATURE:

(D) OTHER INFORMATION: The sequence has an acetylated Nterminus (i.e., $N-\alpha-Ac$), rather than an amino N-terminus (i.e., H2N-). The sequence has an amide C-terminus (i.e., CO-NH2), rather than a carboxyl C-terminus (i.e., CO-OH).

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:8:

Arg His Phe Leu Asn Leu Val Thr Arg Gln Arg Tyr

- (2) INFORMATION FOR SEQ ID NO:9:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 36 amino acids (B) TYPE: amino acid

 - (C) STRANDEDNESS: N/A
 - (D) TOPOLOGY: linear
- (ix) FEATURE:

(D) OTHER INFORMATION: Xaa in position 26 is an abbreviation of im-DNP-His. The sequence has an acetylated Nterminus (i.e., $N-\alpha-Ac$), rather than an amino N-terminus (i.e., H2N-). The sequence has an amide C-terminus (i.e., CO-NH2), rather than a carboxyl C-terminus (i.e., CO-OH).

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:9:

Tyr Pro Ala Lys Pro Glu Ala Pro Gly Glu Asp Ala Ser Pro Glu Glu 5 10 1 Leu Ser Arg Tyr Tyr Ala Ser Leu Arg Xaa Tyr Leu Asn Leu Val Thr 20 Arg Gln Arg Tyr 35

- (2) INFORMATION FOR SEQ ID NO:10:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 15 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: N/A (D) TOPOLOGY: linear
- (ix) FEATURE:

(D) OTHER INFORMATION: The sequence has an amide Cterminus (i.e., CO-NH2), rather than a carboxyl C-terminus (i.e., CO-OH).

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:10:

Ala Ser Leu Arg His Tyr Leu Asn Leu Val Thr Arg Gln Arg Tyr

(2) INFORMATION FOR SEQ ID NO:11:

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- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 15 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: N/A
 - (D) TOPOLOGY: linear
- (ix) FEATURE:

(D) OTHER INFORMATION: The sequence has an amide Cterminus (i.e., CO-NH2), rather than a carboxyl C-terminus (i.e., CO-

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:11:

Ala Ser Leu Arg His Tyr Leu Asn Leu Val Ala Arg Gln Arg Tyr

- (2) INFORMATION FOR SEQ ID NO:12:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 15 amino acids
 - (B) TYPE: amino acid

 - (C) STRANDEDNESS: N/A (D) TOPOLOGY: linear
- (ix) FEATURE:

(D) OTHER INFORMATION: The sequence has an amide Cterminus (i.e., CO-NH2), rather than a carboxyl C-terminus (i.e., CO-OH).

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:12:

Ala Ala Leu Arg His Tyr Leu Asn Leu Val Ala Arg Gln Arg Tyr 10. 1

- (2) INFORMATION FOR SEQ ID NO:13:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 15 amino acids
 - (B) TYPE: amino acid (C) STRANDEDNESS: N/A
 - (D) TOPOLOGY: linear
- (ix) FEATURE:

(D) OTHER INFORMATION: The sequence has an amide Cterminus (i.e., CO-NH2), rather than a carboxyl C-terminus (i.e., CO-OH).

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:13:

Ala Ser Leu Arg His Tyr Glu Asn Leu Val Thr Arg Gln Arg Tyr 1

- (2) INFORMATION FOR SEQ ID NO:14:
- (i) SEQUENCE CHARACTERISTICS:

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- (A) LENGTH: 15 amino acids
- (B) TYPE: amino acid
- (C) STRANDEDNESS: N/A (D) TOPOLOGY: linear

(ix) FEATURE:

(D) OTHER INFORMATION: The sequence has an acetylated Nterminus (i.e., $N-\alpha-Ac$), rather than an amino N-terminus (i.e., H2N-). The sequence has an amide C-terminus (i.e., CO-NH2), rather than a carboxyl C-terminus (i.e., CO-OH).

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:14:

Ala Ser Leu Arg His Tyr Leu Asn Leu Val Thr Arg Gln Arg Tyr 10

- (2) INFORMATION FOR SEQ ID NO:15:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 15 amino acids
 - (B) TYPE: amino acid

 - (C) STRANDEDNESS: N/A (D) TOPOLOGY: linear
- (ix) FEATURE:

(D) OTHER INFORMATION: Xaa in position 5 is an abbreviation of p.Cl.Pro. The sequence has an acetylated N-terminus (i.e., $N-\alpha-Ac$), rather than an amino N-terminus (i.e., H2N-). sequence has an amide C-terminus (i.e., CO-NH2), rather than a carboxyl C-terminus (i.e., CO-OH).

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:15:

Ala Ser Leu Arg Xaa Tyr Leu Asn Leu Val Thr Arg Gln Arg Tyr 1 15

- (2) INFORMATION FOR SEQ ID NO:16:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 15 amino acids (B) TYPE: amino acid (C) STRANDEDNESS: N/A

 - (D) TOPOLOGY: linear
- (ix) FEATURE:

(D) OTHER INFORMATION: The sequence has an acetylated Nterminus (i.e., $N-\alpha-Ac$), rather than an amino N-terminus (i.e., H2N-). The sequence has an amide C-terminus (i.e., CO-NH2), rather than a carboxyl C-terminus (i.e., CO-OH).

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:16:

Ala Ser Leu Arg His Tyr Glu Asn Leu Val Thr Arg Gln Arg Tyr 10

(2) INFORMATION FOR SEQ ID NO:17:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 15 amino acids (B) TYPE: amino acid

 - (C) STRANDEDNESS: N/A
 - (D) TOPOLOGY: linear

(ix) FEATURE:

(D) OTHER INFORMATION: Xaa in position 15 is an abbreviation of N-Me-Tyr. The sequence has an acetylated N-terminus (i.e., $N-\alpha-Ac$), rather than an amino N-terminus (i.e., H2N-). The sequence has an amide C-terminus (i.e., CO-NH2), rather than a carboxyl C-terminus (i.e., CO-OH).

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:17:

Ala Ser Leu Arg His Tyr Glu Asn Leu Val Thr Arg Gln Arg Xaa

- (2) INFORMATION FOR SEQ ID NO:18:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 15 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: N/A (D) TOPOLOGY: linear
- (ix) FEATURE:

(D) OTHER INFORMATION: The sequence has a myristoylated N-terminus (i.e., N- α -myristoyl), rather than an amino N-terminus (i.e., H2N-). The sequence has an amide C-terminus (i.e., CO-NH2), rather than a carboxyl C-terminus (i.e., CO-OH).

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:18:

Ala Ser Leu Arg His Tyr Leu Asn Leu Val Thr Arg Gln Arg Tyr

- (2) INFORMATION FOR SEQ ID NO:19:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 15 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: N/A (D) TOPOLOGY: linear
- (ix) FEATURE:

(D) OTHER INFORMATION: The sequence has a napthalenoacetyl N-terminus (i.e., N- α -napthaleneacetyl), rather than an amino N-terminus (i.e., H2N-). The sequence has an amide Cterminus (i.e., CO-NH2), rather than a carboxyl C-terminus (i.e., CO-OH).

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:19:

Ala Ser Leu Arg His Tyr Leu Asn Leu Val Thr Arg Gln Arg Tyr 10

- (2) INFORMATION FOR SEQ ID NO:20:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 15 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: N/A
 - (D) TOPOLOGY: linear

(ix) FEATURE:

(D) OTHER INFORMATION: The sequence has an acetylated Nterminus (i.e., N- α -Ac), rather than an amino N-terminus (i.e., H2N-). The sequence has an amide C-terminus (i.e., CO-NH2), rather than a carboxyl C-terminus (i.e., CO-OH).

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:20:

Ala Ser Leu Arg His Tyr Leu Asn Leu Val Thr Arg Gln Arg Tyr

- (2) INFORMATION FOR SEQ ID NO:21:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 15 amino acids

 - (B) TYPE: amino acid (C) STRANDEDNESS: N/A
 - (D) TOPOLOGY: linear
- (ix) FEATURE:

(D) OTHER INFORMATION: Xaa in position 6 is an abbreviation of Bth. The sequence has an acetylated N-terminus (i.e., $N-\alpha-Ac$), rather than an amino N-terminus (i.e., H2N-). The sequence has an amide C-terminus (i.e., CO-NH2), rather than a carboxyl C-terminus (i.e., CO-OH).

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:21:

Ala Ser Leu Arg His Xaa Leu Asn Leu Val Thr Arg Gln Arg Tyr 5

- (2) INFORMATION FOR SEQ ID NO:22:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 15 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: N/A
 - (D) TOPOLOGY: linear
- (ix) FEATURE:

(D) OTHER INFORMATION: Xaa in position 6 is an abbreviation of Bip. The sequence has an acetylated N-terminus (i.e., $N-\alpha-Ac$), rather than an amino N-terminus (i.e., H2N-). sequence has an amide C-terminus (i.e., CO-NH2), rather than a carboxyl C-terminus (i.e., CO-OH).

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:22:

Ala Ser Leu Arq His Xaa Leu Asn Leu Val Thr Arg Gln Arg Tyr

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(2) INFORMATION FOR SEQ ID NO:23:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 15 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: N/A
 - (D) TOPOLOGY: linear

(ix) FEATURE:

(D) OTHER INFORMATION: Xaa in position 6 is an abbreviation of Nal. The sequence has an acetylated N-terminus (i.e., N- α -Ac), rather than an amino N-terminus (i.e., H2N-). The sequence has an amide C-terminus (i.e., CO-NH2), rather than a carboxyl C-terminus (i.e., CO-OH).

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:23:

Ala Ser Leu Arg His Xaa Leu Asn Leu Val Thr Arg Gln Arg Tyr
1 5 10 15

(2) INFORMATION FOR SEQ ID NO:24:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 36 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: N/A
 - (D) TOPOLOGY: linear

(ix) FEATURE:

(D) OTHER INFORMATION: The sequence has an amide Cterminus (i.e., CO-NH2), rather than a carboxyl C-terminus (i.e., CO-OH).

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:24:

Tyr Pro Ser Lys Pro Asp Asn Pro Gly Glu Asp Ala Pro Ala Glu Asp 1 5 10 15 Met Ala Arg Tyr Tyr Ser Ala Leu Arg His Tyr Ile Asn Leu Ile Thr $\frac{20}{35}$ Arg Gln Arg Tyr

(2) INFORMATION FOR SEQ ID NO:25:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 15 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: N/A
 - (D) TOPOLOGY: linear

(ix) FEATURE:

(D) OTHER INFORMATION: Xaa in position 6 is an abbreviation of Tic. The sequence has an acetylated N-terminus (i.e., N- α -Ac), rather than an amino N-terminus (i.e., H2N-). The

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sequence has an amide C-terminus (i.e., CO-NH2), rather than a carboxyl C-terminus (i.e., CO-OH).

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:25:

Ala Ser Leu Arg His Xaa Leu Asn Leu Val Thr Arg Gln Arg Tyr 5 10

- (2) INFORMATION FOR SEQ ID NO:26:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 11 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: N/A
 - (D) TOPOLOGY: linear
- (ix) FEATURE:

(D) OTHER INFORMATION: The sequence has an acetylated Nterminus (i.e., $N-\alpha-Ac$), rather than an amino N-terminus (i.e., H2N-). The sequence has an amide C-terminus (i.e., CO-NH2), rather than a carboxyl C-terminus (i.e., CO-OH).

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:26:

His Phe Leu Asn Leu Val Thr Arg Gln Arg Tyr 5

- (2) INFORMATION FOR SEQ ID NO:27:
- (i) SEQUENCE CHARACTERISTICS:
- (A) LENGTH: 15 amino acids
 (B) TYPE: amino acid
- (C) STRANDEDNESS: N/A
 - (D) TOPOLOGY: linear
 - (ix) FEATURE:

(D) OTHER INFORMATION: Xaa in position 15 is an abbreviation of Thi. The sequence has an acetylated N-terminus (i.e., $N-\alpha-Ac$), rather than an amino N-terminus (i.e., H2N-). The sequence has an amide C-terminus (i.e., CO-NH2), rather than a carboxyl C-terminus (i.e., CO-OH).

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:27:

Ala Ser Leu Arg His Phe Leu Asn Leu Val Thr Arg Gln Arg Xaa 5 10

- (2) INFORMATION FOR SEQ ID NO:28:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 15 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: N/A (D) TOPOLOGY: linear
- (ix) FEATURE:

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(D) OTHER INFORMATION: Xaa in position 15 is an abbreviation of Thz. The sequence has an acetylated N-terminus (i.e., $N-\alpha-Ac$), rather than an amino N-terminus (i.e., H2N-). The sequence has an amide C-terminus (i.e., CO-NH2), rather than a carboxyl C-terminus (i.e., CO-OH).

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:28:

Ala Ser Leu Arg Xaa Phe Leu Asn Leu Val Thr Arg Gln Arg Tyr

- (2) INFORMATION FOR SEQ ID NO:29:
- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 15 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: N/A (D) TOPOLOGY: linear
- (ix) FEATURE:

(D) OTHER INFORMATION: Xaa in position 6 is an abbreviation of Pcp. The sequence has an acetylated N-terminus (i.e., $N-\alpha-Ac$), rather than an amino N-terminus (i.e., H2N-). The sequence has an amide C-terminus (i.e., CO-NH2), rather than a carboxyl C-terminus (i.e., CO-OH).

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:29:

Ala Ser Leu Arg His Xaa Leu Asn Leu Val Thr Arg Gln Arg Tyr

- (2) INFORMATION FOR SEQ ID NO:30:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 15 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: N/A (D) TOPOLOGY: linear
 - (ix) FEATURE:

(D) OTHER INFORMATION: The sequence has an acetylated Nterminus (i.e., $N-\alpha-Ac$), rather than an amino N-terminus (i.e., H2N-). The sequence has an amide C-terminus (i.e., CO-NH2), rather than a carboxyl C-terminus (i.e., CO-OH).

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:30:

Phe Ser Leu Arg His Phe Leu Asn Leu Val Thr Arg Gln Arg Tyr

What is claimed is:

1. A compound having the formula:

$$R_1$$
 R_2
 R_2
 R_3
 R_4
 R_2
 R_3
 R_4
 R_5
 R_5
 R_5
 R_6
 R_7
 R_8
 R_8
 R_8
 R_8
 R_9
 R_9

5 wherein

Anb.

10

25

X is Cys or is deleted;

Y is a chain of 0-4 amino acids, inclusive, the C-terminal one of which is bonded to R_3 and R_4 ;

 R_1 is H, C_1-C_{12} alkyl, C_6-C_{18} aryl, C_1-C_{12} acyl, C_7-C_{18} aralkyl, or C_7-C_{18} alkaryl;

 R_2 is H, C_1-C_{12} alkyl, C_6-C_{18} aryl, C_1-C_{12} acyl, C_7-C_{18} aralkyl, or C_7-C_{18} alkaryl;

A²² is an aromatic amino acid, Ala, Aib, Anb, N-Me-Ala, or is deleted;

15 A²³ is Ser, Thr, Ala, Aib, N-Me-Ser, N-Me-Thr, N-Me- Ala, D-Trp, or is deleted;

A²⁴ is Leu, Gly, Ile, Val, Trp, Nle, Nva, Aib,

N-Me-Leu, or is deleted;

 A^{25} is Arg, Lys, homo-Arg, diethyl-homo-Arg, Lys- ϵ - C_{10} NH-R (where R is H, a branched or straight chain C_{1} - C_{10} alkyl group, or an aryl group),
Orn or is deleted;

 A^{26} is Ala, His, Thr, 3-Me-His, 1-Me-His, β -pyrozolylalanine, N-Me-His, Arg, Lys, homo-Arg, diethyl-homo-Arg, Lys- ϵ -NH-R (where R is H, a branched chain or straight chain C_1 - C_{10} alkyl group, or an aryl group), Orn,or is deleted;

A²⁷ is an aromatic amino acid other than Tyr;
A²⁸ is Leu, Val, Trp, Nle, Nva, Aib, Anb, or
N-Me-Leu;

A²⁹ is Asn, Ala, Gln, Gly, Trp, or N-Me-Asn; A³⁰ is Leu, Ile, Val, Trp, Nle, Nva, Aib, Anb, or

N-Me-Leu;

5

- A³¹ is Val, Leu, Ile, Trp, Nle, Nva, Aib, Anb, or N-Me-Val;
- A³² is Thr, Ser, N-Me-Ser, N-Me-Thr, or D-Trp;
- R_3 is H, C_1 - C_{12} alkyl, C_6 - C_{18} aryl, C_1 - C_{12} acyl, C_7 - C_{18} aralkyl, or C_7 - C_{18} alkaryl; and
- R_4 is H, C_1-C_{12} alkyl, C_6-C_{18} aryl, C_1-C_{12} acyl, C_7-C_{18} aralkyl, C_7-C_{18} alkaryl, or a pharmaceutically acceptable salt thereof.
- 10 2. The compound of claim 1, wherein A²⁷ is Phe, Nal, Bip, Pcp, Tic, Trp, Trp, Bth, Thi, or Dip.
 - 3. The compound of claim 1, where Y is $A^{33}-A^{34}-A^{35}-A^{36}$ wherein
- A^{33} is Cys, Arg, Lys, homo-Arg, diethyl-homo-Arg, Lys- ϵ -NH-R (where R is H, a branched or straight chain C_1 - C_{10} alkyl group, or C_6 - C_{18} aryl group), or Orn;
 - A³⁴ is Cys, Gln, Asn, Ala, Gly, N-Me-Gln, Aib, or Anb;
- ${
 m A}^{35}$ is Cys, Arg, Lys, homo-Arg, diethyl-homo-Arg, Lys- ϵ -NH-R (where R is H, a branched or straight chain ${
 m C}_1$ - ${
 m C}_{10}$ alkyl group, or ${
 m C}_6$ - ${
 m C}_{18}$ aryl group), or Orn; and
- A³⁶ is an aromatic amino acid, Cys, or a
 25 pharmaceutically acceptable salt thereof.
 - 4. The compound of claim 3, wherein said compound has the formula:

 $N-\alpha-Ac-Ala-Ser-Leu-Arg-His-Phe-Leu-Asn-Leu-Val-Thr-Arg-Gln-Arg-Tyr-NH_2$ (SEQ. ID. NO. 3), or a pharmaceutically acceptable salt thereof.

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The compound of claim 3, wherein said compound has the formula:

H-Ala-Ser-Leu-Arg-His-Phe-Leu-Asn-Leu-Val-Thr-Arg-Gln-Arg-Tyr-NH₂ (SEQ. ID. NO. 4), or a pharmaceutically 5 acceptable salt thereof.

6. The compound of claim 3, wherein said compound has the formula:

N-α-Ac-Ala-Ser-Leu-Arg-His-Trp-Leu-Asn-Leu-Val-Thr-Arg-Gln-Arg-Tyr-NH2 (SEQ. ID. NO. 5), or a pharmaceutically 10 acceptable salt thereof.

The compound of claim 3, wherein said compound has the formula:

N-α-Ac-Ala-Ser-Leu-Arg-His-Thi-Leu-Asn-Leu-Val-Thr-Arg-Gln-Arg-Tyr-NH2 (SEQ. ID. NO. 6), or a pharmaceutically 15 acceptable salt thereof.

The compound of claim 3, wherein said compound has the formula:

N-α-Ac-Tyr-Ser-Leu-Arg-His-Phe-Leu-Asn-Leu-Val-Thr-Arg-Gln-Arg-Tyr-NH2 (SEQ. ID. NO. 7), or a 20 pharmaceutically acceptable salt thereof.

9. A compound having the formula:

$$R_1$$
 R_2
 R_2
 R_2
 R_3
 R_4
 R_2
 R_4
 R_5
 R_5
 R_5
 R_5
 R_5
 R_5
 R_5
 R_5
 R_7
 R_8
 R_8
 R_8
 R_8
 R_9
 R_9

25 wherein

the N-terminal amino acid is bonded to R_1 and R_2 ; Y is a chain of 0-4 amino acids, inclusive the Cterminal one of which is bonded to R3 and R4;

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 R_1 is H, C_1-C_{12} alkyl, C_6-C_{18} aryl, C_1-C_{12} acyl, C_7-C_{18} aralkyl, or C_7-C_{18} alkaryl; R_2 is H, C_1-C_{12} alkyl, C_6-C_{18} aryl, C_1-C_{12} acyl, C_7-C_{18} aralkyl, or C_7-C_{18} alkaryl; A²⁵ is Arg, Lys, homo-Arg, diethyl-homo-Arg, Lys-5 NH-R (where R is H, a branched or straight ϵ – chain C_1-C_{10} alkyl group, or an aryl group), Orn or is deleted; A²⁶ is Ala, His, Thr, 3-Me-His, β -pyrozolylalanine, N-Me-His, Arg, Lys, homo-10 Arg, diethyl-homo-Arg, Lys- ϵ -NH-R (where R is H, a branched or straight chain C₁-C₁₀ alkyl group, or an aryl group), or Orn; A²⁷ is an aromatic amino acid; A²⁸ is Leu, Val, Trp, Nle, Nva, Aib, Anb, or 15 N-Me-Leu; A²⁹ is Asn, Ala, Gln, Gly, Trp, or N-Me-Asn; A³⁰ is Leu, Ile, Val, Trp, Nle, Nva, Aib, Anb, or N-Me-Leu; 20 A³¹ is Val, Leu, Ile, Trp, Nle, Nva, Aib, Anb, or N-Me-Val; A³² is Thr, Ser, N-Me-Ser, N-Me-Thr, or D-Trp; $\rm R_3$ is H, $\rm C_1-\rm C_{12}$ alkyl, $\rm C_6-\rm C_{18}$ aryl, $\rm C_1-\rm C_{12}$ acyl, $\rm C_7-\rm$ C_{18} aralkyl, or C_7 - C_{18} alkaryl; and R_4 is H, 25 C₁- C_{12} alkyl, C_6-C_{18} aryl, C_1-C_{12} acyl, C_7-C_{18} aralkyl or C_7-C_{18} alkaryl, or a

10. The compound of claim 9, wherein ${\tt A}^{27}$ is Phe, Nal, Bip, Pcp, Tic, Trp, Bth, Thi, or Dip.

pharmaceutically acceptable salt thereof.

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- 11. The compound of claim 9, wherein Y is $A^{33}-A^{34}-A^{35}-A^{36}$ wherein
- A^{33} is Arg, Lys, homo-Arg, diethyl-homo-Arg, Lys- ϵ - NH-R (where R is H, a branched or straight chain C_1 - C_{10} alkyl group, or C_6 - C_{18} aryl group), Cys, or Orn;
 - A³⁴ is Cys, Gln, Asn, Ala, Gly, N-Me-Gln, Alb, or Anb;
- A^{35} is Arg, Lys, homo-Arg, diethyl-homo-Arg, Lys10 ϵ NH-R (where R is H, a branched or straight chain C_1 - C_{10} alkyl group, or C_6 - C_{18} aryl group), Cys, or Orn; and
 - A³⁶ is an aromatic amino acid, Cys, or a pharmaceutically acceptable salt thereof.
- 15 12. The compound of claim 11, wherein said compound has the formula: $N-\alpha-Ac-Arg-His-Phe-Leu-Asn-Leu-Val-Thr-Arg-Gln-Arg-Tyr-NH_2 \ (SEQ. ID. NO. 26), or a pharmaceutically acceptable salt thereof.$
- 20 13. A therapeutic composition capable of decreasing excess intestinal water and electrolyte secretion, said composition comprising a therapeutically effective amount of the compound of claim 1 or claim 9, together with a pharmaceutically acceptable carrier substance.
- 14. A method of decreasing excess intestinal water and electrolyte secretion in a mammal, said method comprising administering to said mammal a therapeutically effective amount of the composition of claim 1 or claim 30 9, together with a pharmaceutically acceptable carrier substance.

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15. A method of regulating cell proliferation in a mammal, said method comprising administering to said mammal a therapeutically effective amount of the composition of claim 1 or claim 9, together with a 5 pharmaceutically acceptable carrier substance.

- 16. A method of augmenting nutrient transport in a mammal, said method comprising administering to said mammal a therapeutically effective amount of the composition of claim 1 or claim 9, together with a pharmaceutically acceptable carrier substance.
- 17. A method or regulating lipolysis in a mammal, said method comprising adminsitering to said mammal a therapeutically effective amount of the composition of claim 1 or claim 9, together with a pharmaceutically acceptable carrier substance.
- 18. A method of regulating blood flow in a mammal, said method comprising adminsitering to said mammal a therapeutically effective amount of the composition of claim 1 or claim 9, together with a 20 pharmaceutically acceptable carrier substance.
- 19. A dimeric compound comprising either two peptides of claim 1 or claim 9, or one peptide of claim 1 or one peptide of claim 9, wherein said dimer is formed by either an amide bond, or a disulfide bridge between 25 said two peptides.

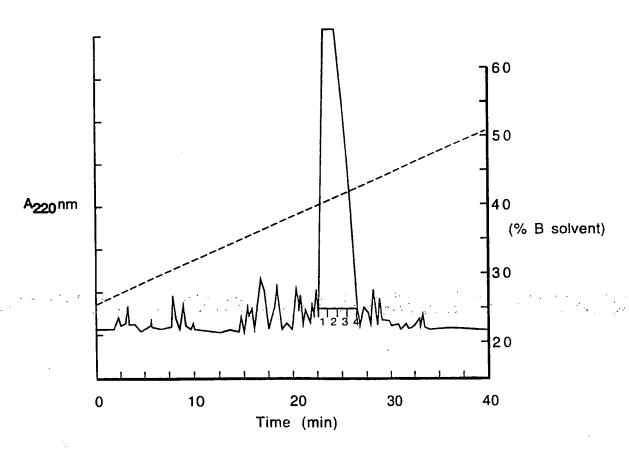


FIG. 1

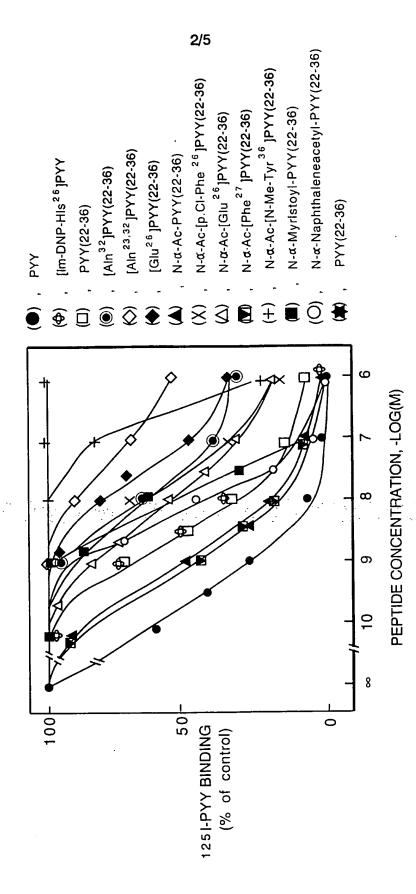


FIG. 2

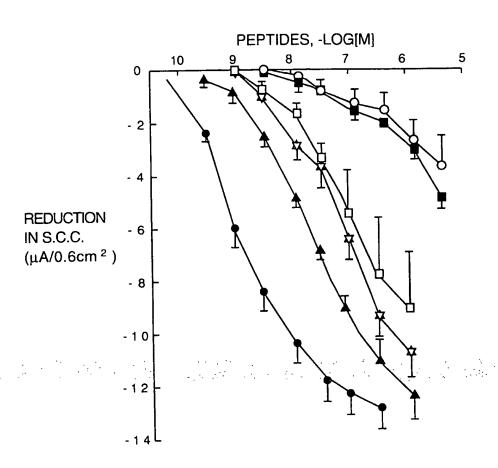


FIG. 3A

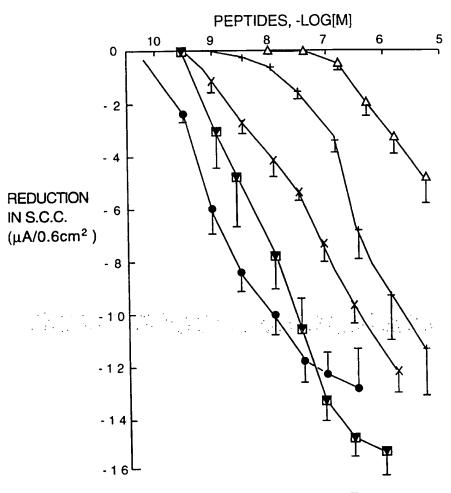
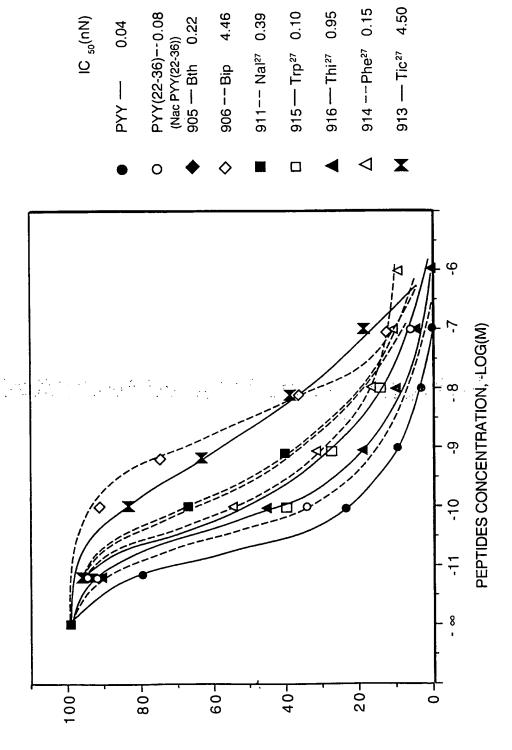


FIG. 3B



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FIG. 4

INTERNATIONAL SEARCH REPORT

International application No. PCT/US96/18374

A. CLASSIFICATION OF SUBJECT MATTER			
IPC(6) :A61K 38/00; C07K 5/00, 7/00, 17/00			
US CL :514/12, 14, 15; 530/324, 326, 327 According to International Patent Classification (IPC) or to both national classification and IPC			
B. FIELDS SEARCHED			
Minimum documentation searched (classification system followed by classification symbols)			
U.S. : 514/12, 14, 15; 530/324, 326, 327.			
Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched			
Electronic data base consulted during the international search (name of data base and, where practicable, search terms used)			
APS, CAS ONLINE, STN, BIOSIS, MEDLINE, EMBASE.			
C. DOCUMENTS CONSIDERED TO BE RELEVANT			
Category*	Citation of document, with indication, where app	propriate, of the relevant passages	Relevant to claim No.
Υ	US 5,328,899 A (BOUBLIK et al.) 12 July 1994, see entire		1-19
T	document.	12 buly 100 if 000 circums	
Y	US 5,026,685 A (BOUBLIK et al.)	1-19	
	document.		
V	TATEMOTO et al. Synthesis of receptor antagonists of 1-19		
Υ	neuropeptide Y. Proc. Natl. Aca		
	1992, Vol 89, pages 1174-1178,		
	17552, Vol. 65, pages 177 175, 555 citation 1875		
Υ	TATEMOTO, K. Neuropeptide Y and Its Receptor		1-19
	Antagonists: Use of an Analog Mixture-Screening Strategy.		
	Annals New York Academy of Sciences. 1990, Vol 611,		
	pages 1-5, see entire document.		
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